<110>   Ekwuribe, Nnochiri
        Radhakrishnan, Balasingam
        Price, Christopher
        Anderson, Wesley
        Ansari, Aslam

<120>   METHODS OF ANALGESIA

<130>   9233-8DV2

<140>   US 09/430,735
<141>   1999-10-29

<150>   US 09/134,803
<151>   1998-08-14

<160>   53

<170>   PatentIn version 3.1

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<220>
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<222>   (6)..(6)
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<400>   1

Tyr Gly Gly Phe Met Lys
1   5

<210>   2
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<220>
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<220>
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<223>   Polymer connected to alpha-amino group

<220>
<221>   MOD_RES
<222>   (6)..(6)
<223>   Polymer connected to epsilon-amino group

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<400>  2

Tyr Gly Gly Phe Met Lys
1               5


<210>  3
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<220>
<223>  Synthetic construct

<220>
<221>  MOD_RES
<222>  (1)..(1)
<223>  Polymer connected to alpha-amino group


<400>  3

Tyr Gly Gly Phe Met Lys
1               5


<210>  4
<211>  6
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<220>
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<220>
<221>  MOD_RES
<222>  (6)..(6)
<223>  AMIDATION


<400>  4

Phe Arg Trp Trp Tyr Lys
1               5


<210>  5
<211>  6
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<220>
<223>  Synthetic construct
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<220>
<221>  MOD_RES
<222>  (1)..(1)
<223>  ACETYLATION


<220>
<221>  MOD_RES
<222>  (6)..(6)
<223>  AMIDATION


<400>  5

Arg Trp Ile Gly Trp Lys
1               5


<210>  6
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<220>
<223>  Synthetic construct

<220>
<221>  MOD_RES
<222>  (6)..(6)
<223>  AMIDATION


<220>
<221>  MISC_FEATURE
<222>  (6)..(6)
<223>  Xaa can be any of the twenty naturally occurring amino acids


<400>  6

Trp Trp Pro Lys His Xaa
1               5


<210>  7
<211>  4
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MOD_RES
<222>  (4)..(4)
<223>  AMIDATION


<220>
<221>  MISC_FEATURE
<222>  (4)..(4)
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<223>   Xaa is either Lys or Arg


<400>   7

Trp Trp Pro Xaa
1


<210>   8
<211>   6
<212>   PRT
<213>   Artificial sequence

<220>
<223>   Synthetic construct

<220>
<221>   MOD_RES
<222>   (6)..(6)
<223>   AMIDATION


<220>
<221>   MISC_FEATURE
<222>   (6)..(6)
<223>   Xaa can be any one of the naturally occurring amino acids


<400>   8

Tyr Pro Phe Gly Phe Xaa
1               5


<210>   9
<211>   7
<212>   PRT
<213>   Artificial sequence

<220>
<223>   Synthetic construct

<220>
<221>   MOD_RES
<222>   (1)..(5)
<223>   Amino acids are in the D-form


<220>
<221>   MOD_RES
<222>   (6)..(6)
<223>   n is 0 or 1


<220>
<221>   MISC_FEATURE
<222>   (7)..(7)
<223>   Xaa is Gly or the D-form of a naturally occurring amino acid
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<220>
<221>  MOD_RES
<222>  (7)..(7)
<223>  Amidation


<400>  9

Ile Met Ser Trp Trp Gly Xaa
1                   5


<210>  10
<211>  6
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
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<222>  (1)..(4)
<223>  Amino acids are in the D-form


<220>
<221>  MISC_FEATURE
<222>  (6)..(6)
<223>  Xaa is Gly or the D-form of a naturally occurring amino acid


<220>
<221>  MOD_RES
<222>  (6)..(6)
<223>  AMIDATION


<400>  10

Ile Met Thr Trp Gly Xaa
1                   5


<210>  11
<211>  4
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is A1, wherein A1 is the D-form of Nve or Mle


<220>
<221>  MISC_FEATURE
<222>  (3)..(3)
```

<223> Xaa is B2, wherein B2 is Gly, Phe, or Trp

<220>
<221> MISC_FEATURE
<222> (4)..(4)
<223> Xaa is C3, wherein C3 is Trp or Nap

<220>
<221> MOD_RES
<222> (4)..(4)
<223> AMIDATION

<400> 11

Tyr Xaa Xaa Xaa
1

<210> 12
<211> 3
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MOD_RES
<222> (1)..(1)
<223> Tyr has at its N-terminus a Me-x-H-y-N group, wherein x is 0, 1,
      or 2; and y is 0, 1, or 2, with the proviso that x and y is never
      greater than 2

<220>
<221> MOD_RES
<222> (1)..(2)
<223> The amine between the first Tyr and the second Tyr is methylated,
      wherein z is 0 or 1

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa is Phe or (D) Phe

<220>
<221> MOD_RES
<222> (3)..(3)
<223> AMIDATION

<400> 12

Tyr Tyr Xaa
1

6

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<210>  13
<211>  6
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (4)..(4)
<223>  Xaa is D4, wherein D4 is Lys or Arg


<220>
<221>  MOD_RES
<222>  (5)..(5)
<223>  His is His-z, wherein z is 0 or 1


<220>
<221>  MISC_FEATURE
<222>  (6)..(6)
<223>  Xaa is Xaa-z, wherein Xaa is a naturally occuring amino acid and
       z is 0 or 1


<220>
<221>  MOD_RES
<222>  (6)..(6)
<223>  AMIDATION


<400>  13

Trp Trp Pro Xaa His Xaa
1                   5


<210>  14
<211>  4
<212>  PRT
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<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<400>  14

Tyr Xaa Phe Phe
1


<210>  15
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<211>    4
<212>    PRT
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<220>
<223>    Synthetic construct

<220>
<221>    MISC_FEATURE
<222>    (2)..(2)
<223>    Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>    MOD_RES
<222>    (4)..(4)
<223>    AMIDATION


<400>    15

Tyr Xaa Phe Phe
1


<210>    16
<211>    4
<212>    PRT
<213>    Artificial sequence

<220>
<223>    Synthetic construct

<220>
<221>    MOD_RES
<222>    (1)..(1)
<223>    Tyr is Tyr(N-alpha-Me), i.e. N-alpha-methyl tyrosine


<220>
<221>    MISC_FEATURE
<222>    (2)..(2)
<223>    Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<400>    16

Tyr Xaa Phe Phe
1


<210>    17
<211>    4
<212>    PRT
<213>    Artificial sequence

<220>
<223>    Synthetic construct

<220>
<221>    MOD_RES
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<222>    (1)..(1)
<223>    Tyr is Tyr(N-alpha-Cmp), i.e. N-alpha-cyclopropylmethyltyrosine


<220>
<221>    MISC_FEATURE
<222>    (2)..(2)
<223>    Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<400>    17

Tyr Xaa Phe Phe
1


<210>    18
<211>    4
<212>    PRT
<213>    Artificial sequence

<220>
<223>    Synthetic construct

<220>
<221>    MOD_RES
<222>    (1)..(1)
<223>    Tyr is Tyr(N-alpha-hex), i.e. N-alpha-hexyltyrosine


<220>
<221>    MISC_FEATURE
<222>    (2)..(2)
<223>    Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<400>    18

Tyr Xaa Phe Phe
1


<210>    19
<211>    4
<212>    PRT
<213>    Artificial sequence

<220>
<223>    Synthetic construct

<220>
<221>    MOD_RES
<222>    (1)..(1)
<223>    Tyr is Tyr(N-alpha-Et2), i.e. N-alpa-diethyltyrosine


<220>
<221>    MISC_FEATURE
<222>    (2)..(2)
<223>    Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
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<400> 19

Tyr Xaa Phe Phe
1

<210> 20
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MOD_RES
<222> (1)..(1)
<223> Tyr is Dmt, i.e. 2,6-dimethyltyrosine

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<400> 20

Tyr Xaa Phe Phe
1

<210> 21
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
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<223> Tyr is Dmt, i.e. 2,6-dimethyltyrosine

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<220>
<221> MOD_RES
<222> (4)..(4)
<223> AMIDATION

<400> 21

Tyr Xaa Phe Phe
1

<210> 22
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MOD_RES
<222> (1)..(1)
<223> Tyr is H-Tyr(3-F), i.e. 3-fluorotyrosine

<220>
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<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<400> 22

Tyr Xaa Phe Phe
1

<210> 23
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
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<223> Tyr is H-Tyr(3-Cl), i.e. 3-chlorotyrosine

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<400> 23

Tyr Xaa Phe Phe
1

<210> 24
<211> 4
<212> PRT
<213> Artificial sequence

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<220>
<223>  Synthetic construct

<220>
<221>  MOD_RES
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<223>  Tyr is H-Tyr (3-Br), i.e. 3-bromotyrosine


<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<400>  24

Tyr Xaa Phe Phe
1


<210>  25
<211>  4
<212>  PRT
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<220>
<223>  Synthetic construct

<220>
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<223>  Tyr is Dmt, i.e. 2,6-dimethyltyrosine


<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic-psi-[CH2-], i.e. 3-methyl-1,2,3,4-tetrahydroisoquinoli
        ne-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (2)..(3)
<223>  nonpetidyl bond


<400>  25

Tyr Xaa Phe Phe
1


<210>  26
<211>  4
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct
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<220>
<221>  MOD_RES
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<223>  Tyr is Dmt, i.e. 2,6-dimethyltyrosine


<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic-psi-[CH2-], i.e. 3-methyl-1,2,3,4-tetrahydroisoquinoli
       ne-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (2)..(3)
<223>  nonpeptidyl bond


<220>
<221>  MOD_RES
<222>  (4)..(4)
<223>  AMIDATION


<400>  26

Tyr Xaa Phe Phe
1


<210>  27
<211>  4
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic-psi-[CH2-], i.e. 3-methyl-1,2,3,4-tetrahydroisoquinoli
       ne-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (3)..(3)
<223>  Phe is -NCH3]Phe, i.e. N-methylphenylalanine


<400>  27

Tyr Xaa Phe Phe
1


<210>  28
<211>  4
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<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic-psi-[CH2-], i.e. 3-methyl-1,2,3,4-tetrahydroisoquinoli
       ne-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (3)..(3)
<223>  Phe is -NH]Hfe, i.e. homophenylalanine


<400>  28

Tyr Xaa Phe Phe
1


<210>  29
<211>  4
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MOD_RES
<222>  (1)..(1)
<223>  Tyr is Tyr(NMe), i.e. N-methyltyrosine


<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic-psi-[CH2-], i.e. 3-methyl-1,2,3,4-tetrahydroisoquinoli
       ne-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (3)..(3)
<223>  Phe is -NH]Hfe, i.e. homophenylalanine


<400>  29

Tyr Xaa Phe Phe
1


<210>  30
<211>  4
<212>  PRT
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<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<220>
<221> MOD_RES
<222> (3)..(3)
<223> Gly is Phg, i.e. phenylglycine

<400> 30

Tyr Xaa Gly Phe
1

<210> 31
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
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<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<400> 31

Tyr Xaa Trp Phe
1

<210> 32
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
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<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<220>
<221> MOD_RES
<222> (4)..(4)
<223> AMIDATION

<400> 32

Tyr Xaa Trp Phe
1

<210> 33
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MISC_FEATURE
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<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<400> 33

Tyr Xaa His Phe
1

<210> 34
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<212> PRT
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<220>
<223> Synthetic construct

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid

<220>
<221> MOD_RES
<222> (3)..(3)
<223> Ala is 2-Nal, i.e. 3-(2'-napthyl)alanine

<400> 34

Tyr Xaa Ala Phe
1

<210> 35
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

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<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>  MISC_FEATURE
<222>  (3)..(3)
<223>  Xaa is Atc, i.e. 2-aminotetralin-2-carboxylic acid


<400>  35

Tyr Xaa Xaa Phe
1


<210>  36
<211>  4
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (4)..(4)
<223>  Phe is Phe(pNO2), i.e. 4-nitrophenylalanine


<400>  36

Tyr Xaa Phe Phe
1


<210>  37
<211>  4
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>  MOD_RES
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<222>    (4)..(4)
<223>    Phe is Phe(pNO2), i.e. 4-nitrophenylalanine


<400>    37

Tyr Xaa Trp Phe
1


<210>    38
<211>    4
<212>    PRT
<213>    Artificial sequence

<220>
<223>    Synthetic construct

<220>
<221>    MISC_FEATURE
<222>    (2)..(2)
<223>    Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>    MOD_RES
<222>    (4)..(4)
<223>    AMIDATION


<400>    38

Tyr Xaa Phe Trp
1


<210>    39
<211>    7
<212>    PRT
<213>    Artificial sequence

<220>
<223>    Synthetic construct

<220>
<221>    MISC_FEATURE
<222>    (2)..(2)
<223>    Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>    MOD_RES
<222>    (7)..(7)
<223>    AMIDATION


<400>    39

Tyr Xaa Phe Phe Val Val Gly
1                    5
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<210>  40
<211>  7
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (7)..(7)
<223>  AMIDATION


<400>  40

Tyr Xaa Phe Phe Tyr Pro Ser
1               5


<210>  41
<211>  7
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (7)..(7)
<223>  AMIDATION


<400>  41

Tyr Xaa Trp Phe Tyr Pro Ser
1               5


<210>  42
<211>  7
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct
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<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (4)..(4)
<223>  Phe is Phe(pNO2), i.e.  4-nitrophenylalanine


<220>
<221>  MOD_RES
<222>  (7)..(7)
<223>  AMIDATION


<400>  42

Tyr Xaa Trp Phe Tyr Pro Ser
1                 5


<210>  43
<211>  7
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
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<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (6)..(6)
<223>  Nle


<220>
<221>  MOD_RES
<222>  (7)..(7)
<223>  AMIDATION


<400>  43

Tyr Xaa Phe Phe Leu Leu Asp
1                 5


<210>  44
<211>  3
<212>  PRT
<213>  Artificial sequence
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<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<400>  44

Tyr Xaa Phe
1


<210>  45
<211>  3
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
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<223>  Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid


<220>
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<222>  (3)..(3)
<223>  AMIDATION


<400>  45

Tyr Xaa Phe
1


<210>  46
<211>  3
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa is Tic-psi-[CH2-], i.e. 3-methyl-1,2,3,4-tetrahydroisoquinoli
       ne-3-carboxylic acid


<220>
<221>  MOD_RES
<222>  (2)..(3)
<223>  nonpeptidyl bond
```

<400> 46

Tyr Xaa Phe
1

<210> 47
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic-psi-[CH2-], i.e. 3-methyl-1,2,3,4-tetrahydroisoquinoli
      ne-3-carboxylic acid

<220>
<221> MOD_RES
<222> (2)..(3)
<223> nonpeptidyl bond

<400> 47

Tyr Xaa Phe Phe
1

<210> 48
<211> 5
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<400> 48

Tyr Gly Gly Phe Met
1   5

<210> 49
<211> 6
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<400> 49

Tyr Gly Gly Phe Met Lys
1   5

<210> 50
<211> 6
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MOD_RES
<222> (1)..(1)
<223> NH2 of Tyr is blocked by butyloxycarbonyl group

<400> 50

Tyr Gly Gly Phe Leu Lys
1   5

<210> 51
<211> 6
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MOD_RES
<222> (1)..(1)
<223> NH2 of Tyr is blocked by butyloxycarbonyl group

<400> 51

Tyr Gly Gly Phe Leu Lys
1   5

<210> 52
<211> 6
<212> PRT
<213> Artificial sequence

<220>
<223> Synthetic construct

<220>
<221> MOD_RES
<222> (1)..(1)
<223> NH2 of Tyr is blocked by butyloxycarbonyl group

<220>
<221> MOD_RES
<222> (6)..(6)
<223> Polymer connected to epsilon-amino group

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<400>  52

Tyr Gly Gly Phe Leu Lys
1                   5


<210>  53
<211>  2
<212>  PRT
<213>  Artificial sequence

<220>
<223>  Synthetic construct

<220>
<221>  MOD_RES
<222>  (1)..(1)
<223>  Tyr has at its N-terminus a Me-x-H-y-N group, wherein x is 0, 1,
       or 2; and y is 0, 1, or 2, with the proviso that x and y is never
       greater than 2


<220>
<221>  MOD_RES
<222>  (1)..(2)
<223>  The amine between the first Tyr and the second Tyr is methylated,
       wherein z is 0 or 1


<220>
<221>  MOD_RES
<222>  (2)..(2)
<223>  AMIDATION


<220>
<221>  MOD_RES
<222>  (2)..(2)
<223>  The amidated C-terminal is benzylated (NHBzl)


<400>  53

Tyr Tyr
1
```